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**ENTERED** OIPE**RAW SEQUENCE LISTING**

PATENT APPLICATION: US/09/936,145

DATE: 03/15/2002

TIME: 14:44:04

Input Set : A:\011309.txt

Output Set: N:\CRF3\03152002\I936145.raw

#9

3 <110> APPLICANT: Inoue, Yasushi
 4 Fushimi, Naoya
 5 Mizubuchi, Hiroyuki
 6 Yamamoto, Yoshie
 7 Ohshima, Yoshie
 8 Yasutake, Nozomu
 9 Miyoshi, Shinsuke
 11 <120> TITLE OF INVENTION: Promoters
 13 <130> FILE REFERENCE: 3274-011309
 15 <140> CURRENT APPLICATION NUMBER: 09/936,145
 16 <141> CURRENT FILING DATE: 2001-09-07
 18 <150> PRIOR APPLICATION NUMBER: PCT/JP00/01415
 19 <151> PRIOR FILING DATE: 2000-03-08
 21 <150> PRIOR APPLICATION NUMBER: US11/060904
 22 <151> PRIOR FILING DATE: 1999-03-08
 24 <150> PRIOR APPLICATION NUMBER: US11/286034
 25 <151> PRIOR FILING DATE: 1999-10-06
 27 <160> NUMBER OF SEQ ID NOS: 22
 29 <170> SOFTWARE: Microsoft Word 97 SR-2
 31 <210> SEQ ID NO: 1
 32 <211> LENGTH: 249
 33 <212> TYPE: DNA
 34 <213> ORGANISM: Bacillus amyloliquefaciens
 36 <400> SEQUENCE: 1
 37 gccgcgcaca tacgaaaaga ctggctgaaa acattgagcc tttgatgact gatgatttgg 60
 38 ctgaagaagt ggatcgattg tttgagaaaa gaagaagacc ataaaaatac cttgtctgtc 120
 39 atcagacagg gtatttttta tgctgtccag actgtccgct gtgtaaaaaa taggaataaa 180
 40 ggggggttgt tattatttta ctgatatgta aaatataatt tgtataagaa aatgagaggg 240
 41 agaggatcc 249
 43 <210> SEQ ID NO: 2
 44 <211> LENGTH: 270
 45 <212> TYPE: DNA
 46 <213> ORGANISM: Bacillus amyloliquefaciens
 48 <400> SEQUENCE: 2
 49 gccccgcaca tacgaaaaga ctggctgaaa acattgagcc tttgatgact gatgatttgg 60
 50 ctgaagaagt ggatcgattg tttgagaaaa gaagaagacc ataaaaatac cttgtctgtc 120
 51 atcagacagg gtatttttta tgctgtccag actgtccgct gtgtaaaaaa taggaataaa 180
 52 ggggggttgt tattatttta ctgatatgta aaatataatt tgtataagaa aatgagaggg 240
 53 agaggatcc ccgggtaccga gtcgaattc 270
 55 <210> SEQ ID NO: 3
 56 <211> LENGTH: 29
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Artificial Sequence

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60 <220> FEATURE:
61 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
63 <400> SEQUENCE: 3
64 cgctctagag ccccgccacat acgaaaaga 29
66 <210> SEQ ID NO: 4
67 <211> LENGTH: 35
68 <212> TYPE: DNA
69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of a primer
73 for introducing a restriction site
75 <400> SEQUENCE: 4
76 cgcgaattcg gatcctctcc ctctcatttt cttat 35
78 <210> SEQ ID NO: 5
79 <211> LENGTH: 50
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of a primer
85 for introducing a restriction site
87 <400> SEQUENCE: 5
88 cgcgaattcg agctcggtac ccgggggatcc tctccctctc attttcttat 50
90 <210> SEQ ID NO: 6
91 <211> LENGTH: 29
92 <212> TYPE: DNA
93 <213> ORGANISM: Artificial Sequence
95 <220> FEATURE:
96 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
98 <400> SEQUENCE: 6
99 cgcggatcca tgtattacaa caggttggt 29
101 <210> SEQ ID NO: 7
102 <211> LENGTH: 29
103 <212> TYPE: DNA
104 <213> ORGANISM: Artificial Sequence
106 <220> FEATURE:
107 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
109 <400> SEQUENCE: 7
110 cgcgaattct cacacatact ccttcgtat 29
112 <210> SEQ ID NO: 8
113 <211> LENGTH: 29
114 <212> TYPE: DNA
115 <213> ORGANISM: Artificial Sequence
117 <220> FEATURE:
118 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
120 <400> SEQUENCE: 8
121 cgcggatcca tgtcttggtc aattagctc 29
123 <210> SEQ ID NO: 9
124 <211> LENGTH: 29
125 <212> TYPE: DNA

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126 <213> ORGANISM: Artificial Sequence
128 <220> FEATURE:
129 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
131 <400> SEQUENCE: 9
132 aaagaattct taatcaacac gcccgttat 29
134 <210> SEQ ID NO: 10
135 <211> LENGTH: 26
136 <212> TYPE: DNA
137 <213> ORGANISM: Artificial Sequence
139 <220> FEATURE:
140 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
142 <400> SEQUENCE: 10
143 gtttcctctc cctctcattt tcttat 26
145 <210> SEQ ID NO: 11
146 <211> LENGTH: 20
147 <212> TYPE: DNA
148 <213> ORGANISM: Artificial Sequence
150 <220> FEATURE:
151 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
153 <400> SEQUENCE: 11
154 atgtattaca acaggttggt 20
156 <210> SEQ ID NO: 12
157 <211> LENGTH: 20
158 <212> TYPE: DNA
159 <213> ORGANISM: Artificial Sequence
161 <220> FEATURE:
162 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
164 <400> SEQUENCE: 12
165 atgtcttggt caattagctc 20
167 <210> SEQ ID NO: 13
168 <211> LENGTH: 29
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
175 <400> SEQUENCE: 13
176 cgcgaattca tgtattacaa caggttggt 29
178 <210> SEQ ID NO: 14
179 <211> LENGTH: 29
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
186 <400> SEQUENCE: 14
187 cgcgaattca tgtcttggtc aattagctc 29
189 <210> SEQ ID NO: 15
190 <211> LENGTH: 1581
191 <212> TYPE: DNA
192 <213> ORGANISM: Agrobacterium radiobacter M36

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194 <220> FEATURE:
195 <221> NAME/KEY: Promoter
196 <222> LOCATION: 314..316
198 <220> FEATURE:
199 <221> NAME/KEY: Terminator
200 <222> LOCATION: 1559..1561
202 <220> FEATURE:
203 <221> NAME/KEY: Gene
204 <222> LOCATION: 341..1558
205 <223> OTHER INFORMATION: MIase structural gene
207 <400> SEQUENCE: 15
208 gatctgcgtg cccatggcac cgtcgagaat gaggatgcgt tcgctggcag cctcgcgcag 60
209 cgccttgaaa atttcgcgc cgtcgcgctt tgccccttca gggccaaaca gatcgtcaaa 120
210 cacgggcaca ctctcattt cgatttgcaa gatcgcaagt cgtcaagtca cataaagata 180
211 tgtttatgtc aatatactt caagggacag gcatggcttt gcgtcgttgc gtcacgttac 240
212 gaaatatcgc tgacagatga caggtttata cggcaaggat ataagccgaa gcagcaaacy 300
213 catggaggac gcaatgcccg aagacgatca caacagccgc aactggaata cctgcacctg 360
214 gcaccgccag tggctggtga aacaggccga gggacttttc gacttcttcc agtatcgcgc 420
215 cctcaatccc gccggcggtt tcttcgatct cgacgccaa ggcgcgcgcgc tgcaggcaaa 480
216 cgatcccgct gcgggcatcc atgcctctgc gcgcattggt cattgcttct ccatcgccca 540
217 cctgctcggc cggcggggct ggcgcgatat cgtcgaccac ggcattgacct atctctggaa 600
218 caaacaccgc gatggcgaa atggcggtta tttctggcag gtgaacgatg ccggcccagt 660
219 ggacgccacc aagcagggtt atggccacgc ctctcgtgct ctggccgcct ctccgcgcaa 720
220 gaccgtcggc caccgcgtgg ccgaccggat gctggctgat attaccgaag tgctggaaag 780
221 tcgtttctgg gaagaaaaac atggcgccat cgccgaggaa ttcaatcgcg actggtcgcc 840
222 catcgacaat tatcgcgccc agaactccaa tatgcacct accgaagcgc tgatggccgc 900
223 ctatgaggtg accggcgaca ataactatct cagcaaggcc gaacgcattc ccatctcgt 960
224 catccgtcgc cgcgcgcgcg agctggattt ccgcgtgccc gagcatttcg acgacaactg 1020
225 gacgtcggac aaggactatc gcggcaacga aatgttccgc ccctccggct ccacccccgg 1080
226 ccaactggct gaattggcgc gtctcattct gcaattgttg aactgggccc aacgcgcgca 1140
227 cgactggatg ccggtcgcgc ccaaattcct ctctcgtcag tccatggcgc tgggctggga 1200
228 ccgggaaaag ggcggcttct tttatacgt ggactggaat gacaatccc acaagcgggc 1260
229 aaagctctgg tggcccatgt ccgaggcggc gggcgcggcc catttctca acgagaacct 1320
230 gccggcggtt ggcttctacg aagacagcta tcgtcgcata tggagcacca tcgccaaaca 1380
231 ctccatcgac catgccaatg gcggctggca tgaggaaact acggaagatc tggttccgc 1440
232 ccacacgcta ttcccaggca agggcgatat ctaccatgcg ctccaggcct gcctcatccc 1500
233 gcttttcccg gcgacgggca gcctgacgaa ggtgatcaag gaaagcggcg gggattatta 1560
234 aggcgtctg cggccaatag c 1581
236 <210> SEQ ID NO: 16
237 <211> LENGTH: 39
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
244 <400> SEQUENCE: 16
245 gcatctcgag catatgcgga tcctctccct ctcatcttc 39
247 <210> SEQ ID NO: 17
248 <211> LENGTH: 31
249 <212> TYPE: DNA

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250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
255 <400> SEQUENCE: 17
256 gcatctcgag ggtaataaaa aaacacctcc a 31
258 <210> SEQ ID NO: 18
259 <211> LENGTH: 30
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
266 <400> SEQUENCE: 18
267 gcatgaattc aaagcagcga tcccgatgaa 30
269 <210> SEQ ID NO: 19
270 <211> LENGTH: 283
271 <212> TYPE: DNA
272 <213> ORGANISM: Bacillus amyloliquefaciens
274 <400> SEQUENCE: 19
275 ctcgagggta ataaaaaac acctccaagc tgagtgcggg taccagcttg gaggtgcgtt 60
276 ttttttttca gccgtatgac aaggctcgga tcaggtgtga caataacggt atgctggctg 120
277 tcataggtga caaatccggg ttttgcgcg tttggctttt tcacatgtct gatttttgta 180
278 taatcaacag gcaaggagcc ggaatctttc gcottggaaa aataagcggc gatcgtagct 240
279 gcttccaata tggattgttc atcgggatcg ctgctttgaa ttc 283
281 <210> SEQ ID NO: 20
282 <211> LENGTH: 28
283 <212> TYPE: DNA
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
289 <400> SEQUENCE: 20
290 gcatcatatg cccgaagacg atcacaac 28
292 <210> SEQ ID NO: 21
293 <211> LENGTH: 31
294 <212> TYPE: DNA
295 <213> ORGANISM: Artificial Sequence
297 <220> FEATURE:
298 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
300 <400> SEQUENCE: 21
301 gcatctcgag ttaataatcc ccgccgcttt c 31
303 <210> SEQ ID NO: 22
304 <211> LENGTH: 21
305 <212> TYPE: DNA
306 <213> ORGANISM: Artificial Sequence
308 <220> FEATURE:
309 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
311 <400> SEQUENCE: 22
312 atgcccgaag acgatcaca c 21

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/936,145

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TIME: 14:44:06

Input Set : A:\011309.txt

Output Set: N:\CRF3\03152002\I936145.raw